

ABSTRACT

The present application provides polynucleic acid sequences of 8 or more contiguous nucleotides selected from an HCV subtype 3c genomic sequence selected from the region spanning positions 1 to 957 of the Core of Core/E1 region of HCV subtype 3c, wherein said polynucleic acid sequence is capable of hybridizing to HCV type 3c, but not another type or subtype of HCV; or the complement of the polynucleic acid, wherein the polynucleic acid contains at least one genotype-specific nucleotide. Methods and means of using and making the described sequences are also provided.